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#2

OIPE

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/997,003

DATE: 08/09/2002 p.6
TIME: 14:56:28

Input Set : A:\PA003P1.seqList.txt
Output Set: N:\CRF4\08092002\I997003.raw

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      1 <110> APPLICANT: Rosen et al.
      3 <120> TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
      5 <130> FILE REFERENCE: PA003P1
C-->  7 <140> CURRENT APPLICATION NUMBER: US/09/997,003
C-->  7 <141> CURRENT FILING DATE: 2001-11-30
      7 <150> PRIOR APPLICATION NUMBER: unassigned
      8 <151> PRIOR FILING DATE: 2001-11-30
     10 <150> PRIOR APPLICATION NUMBER: PCT/US00/22157
     11 <151> PRIOR FILING DATE: 2000-08-11
     13 <150> PRIOR APPLICATION NUMBER: 60/148,680
     14 <151> PRIOR FILING DATE: 1999-08-13
     16 <160> NUMBER OF SEQ ID NOS: 56
     18 <170> SOFTWARE: PatentIn Ver. 2.0
     20 <210> SEQ ID NO: 1
     21 <211> LENGTH: 733
     22 <212> TYPE: DNA
     23 <213> ORGANISM: Homo sapiens
     25 <400> SEQUENCE: 1
     26 gggatccgga gcccaaatct tctgacaaaa ctcacacatg cccaccgtgc ccagcacctg      60
     27 aattcgaggg tgcaccgtca gtcttcctct tccccccaaa acccaaggac accctcatga     120
     28 tctcccggac tcctgaggtc acatgcgtgg tggtggacgt aagccacgaa gaccctgagg     180
     29 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcataa tgccaagaca aagccgcggg     240
     30 aggagcagta caacagcacg taccgtgtgg tcagcgtcct caccgtcctg caccaggact     300
     31 ggctgaatgg caaggagtac aagtgcaagg tctccaacaa agccctccca acccccatcg     360
     32 agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc     420
     33 catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct     480
     34 atccaagcga catcgccgtg gagtgggaga gcaatgggca gccggagaac aactacaaga     540
     35 ccacgcctcc cgtgctggac tccgacggct ccttcttcct ctacagcaag ctcaccgtgg     600
     36 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc     660
     37 acaaccacta cacgcagaag agcctctccc tgtctccggg taaatgagtg cgacggccgc     720
     38 gactctagag gat                                                         733
     40 <210> SEQ ID NO: 2
     41 <211> LENGTH: 5
     42 <212> TYPE: PRT
     43 <213> ORGANISM: Homo sapiens
     45 <220> FEATURE:
     46 <221> NAME/KEY: Site
     47 <222> LOCATION: (3)
     48 <223> OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
     50 <400> SEQUENCE: 2
W--> 51 Trp Ser Xaa Trp Ser
     52    1               5
     54 <210> SEQ ID NO: 3
```

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Input Set : A:\PA003P1.seqList.txt
Output Set: N:\CRF4\08092002\I997003.raw

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    55 <211> LENGTH: 86
    56 <212> TYPE: DNA
    57 <213> ORGANISM: Artificial Sequence
    59 <220> FEATURE:
    60 <221> NAME/KEY: Primer_Bind
    61 <223> OTHER INFORMATION: Synthetic sequence with 4 tandem copies of the GAS binding
site
    62       found in the IRF1 promoter (Rothman et al., Immunity 1:457-468
    63       (1994)), 18 nucleotides complementary to the SV40 early promoter,
    64       and a Xho I restriction site.
    66 <400> SEQUENCE: 3
    67 gcgcctcgag atttccccga aatctagatt tccccgaaat gatttccccg aaatgatttc    60
    68 cccgaaatat ctgccatctc aattag                                        86
    70 <210> SEQ ID NO: 4
    71 <211> LENGTH: 27
    72 <212> TYPE: DNA
    73 <213> ORGANISM: Artificial Sequence
    75 <220> FEATURE:
    76 <221> NAME/KEY: Primer_Bind
    77 <223> OTHER INFORMATION: Synthetic sequence complementary to the SV40 promter;
includes a
    78       Hind III restriction site.
    80 <400> SEQUENCE: 4
    81 gcggcaagct ttttgcaaag cctaggc                                       27
    83 <210> SEQ ID NO: 5
    84 <211> LENGTH: 271
    85 <212> TYPE: DNA
    86 <213> ORGANISM: Artificial Sequence
    88 <220> FEATURE:
    89 <221> NAME/KEY: Protein_Bind
    90 <223> OTHER INFORMATION: Synthetic promoter for use in biological assays; includes GAS
    91       binding sites found in the IRF1 promoter (Rothman et al., Immunity
    92       1:457-468 (1994)).
    94 <400> SEQUENCE: 5
    95 ctcgagattt ccccgaaatc tagatttccc cgaaatgatt tccccgaaat gatttccccg    60
    96 aaatatctgc catctcaatt agtcagcaac catagtcccg cccctaactc cgcccatccc   120
    97 gcccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa tttttttat    180
    98 ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt   240
    99 ttttggaggc ctaggctttt gcaaaaagct t                                  271
   101 <210> SEQ ID NO: 6
   102 <211> LENGTH: 32
   103 <212> TYPE: DNA
   104 <213> ORGANISM: Artificial Sequence
   106 <220> FEATURE:
   107 <221> NAME/KEY: Primer_Bind
   108 <223> OTHER INFORMATION: Synthetic primer complementary to human genomic EGR-1
promoter
   109       sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); includes a
   110       Xho I restriction site.
   112 <400> SEQUENCE: 6
   113 gcgctcgagg gatgacagcg atagaacccc gg                                 32
   115 <210> SEQ ID NO: 7
```

RAW SEQUENCE LISTING   DATE: 08/09/2002
PATENT APPLICATION: US/09/997,003   TIME: 14:56:28

Input Set : **A:\PA003Pl.seqList.txt**
Output Set: **N:\CRF4\08092002\I997003.raw**

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116 <211> LENGTH: 31
117 <212> TYPE: DNA
118 <213> ORGANISM: Artificial Sequence
120 <220> FEATURE:
121 <221> NAME/KEY: Primer_Bind
122 <223> OTHER INFORMATION: Synthetic primer complementary to human genomic EGR-1
promoter
123        sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); includes a
124        Hind III restriction site.
126 <400> SEQUENCE: 7
127 gcgaagcttc gcgactcccc ggatccgcct c                            31
129 <210> SEQ ID NO: 8
130 <211> LENGTH: 12
131 <212> TYPE: DNA
132 <213> ORGANISM: Homo sapiens
134 <400> SEQUENCE: 8
135 ggggactttc cc                                                 12
137 <210> SEQ ID NO: 9
138 <211> LENGTH: 73
139 <212> TYPE: DNA
140 <213> ORGANISM: Artificial Sequence
142 <220> FEATURE:
143 <221> NAME/KEY: Primer_Bind
144 <223> OTHER INFORMATION: Synthetic primer with 4 tandem copies of the NF-KB binding
site
145        (GGGGACTTTCCC), 18 nucleotides complementary to the 5' end of the
146        SV40 early promoter sequence, and a XhoI restriction site.
148 <400> SEQUENCE: 9
149 gcggcctcga ggggactttc ccggggactt tccggggact ttccgggact ttccatcctg   60
150 ccatctcaat tag                                                73
152 <210> SEQ ID NO: 10
153 <211> LENGTH: 256
154 <212> TYPE: DNA
155 <213> ORGANISM: Artificial Sequence
157 <220> FEATURE:
158 <221> NAME/KEY: Protein_Bind
159 <223> OTHER INFORMATION: Synthetic promoter for use in biological assays; includes
NF-KB
160        binding sites.
162 <400> SEQUENCE: 10
163 ctcgagggga ctttcccggg gactttccgg ggactttccg ggactttcca tctgccatct   60
164 caattagtca gcaaccatag tcccgcccct aactccgccc atcccgcccc taactccgcc  120
165 cagttccgcc cattctccgc cccatggctg actaattttt tttatttatg cagaggccga  180
166 ggccgcctcg gcctctgagc tattccagaa gtagtgagga ggctttttg gaggcctagg  240
167 cttttgcaaa aagctt                                             256
169 <210> SEQ ID NO: 11
170 <211> LENGTH: 800
171 <212> TYPE: DNA
172 <213> ORGANISM: Homo sapiens
174 <400> SEQUENCE: 11
175 tcgacccacg cgtccgaact cagactcagc caacagagat tgttgatttg cctcttaagc   60
176 aagagattca ttgcagctca gcatggctca gaccagctca tacttcatgc tgatctcctg  120
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Input Set : **A:\PA003P1.seqList.txt**
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177 cctgatgttt ctgtctcaga gccaaggcca agaggcccag acagagttgc cccaggcccg   180
178 gatcagctgc ccagaaggca ccaatgccta tcgctcctac tgctactact ttaatgaaga   240
179 ccgtgagacc tgggttgatg cagatctcta ttgccagaac atgaattcgg gcaacctggt   300
180 gtctgtgctc acccaggccg agggtgcctt tgtggcctca ctgattaagg agagtggcac   360
181 tgatgacttc aatgtctgga ttggcctcca tgaccccaaa aagaaccgcc gctggcactg   420
182 gagcagtggg tccctggtct cctacaagtc ctggggcatt ggagccccaa gcagtgttaa   480
183 tcctggctac tgtgtgagcc tgacctcaag cacaggattc cagaaatgga aggatgtgcc   540
184 ttgtgaagac aagttctcct ttgtctgcaa gttcaaaaac tagaggcagc tggaaaatac   600
185 atgtctagaa ctgatccagc aattacaacg gagtcaaaaa ttaaaccgga ccatctctcc   660
186 aactcaactc aacctggaca ctctcttctc tgctgagttt gccttgttaa tcttcaatag   720
187 ttttacctac cccagtcttt ggaaccctaa ataataaaaa taaacatgtt tccactatta   780
188 aaacaaaaaa aaaaaaaaaa                                                 800
190 <210> SEQ ID NO: 12
191 <211> LENGTH: 514
192 <212> TYPE: DNA
193 <213> ORGANISM: Homo sapiens
195 <400> SEQUENCE: 12
196 ccacgcgtcc gctgcactct caggtattcc ctgctcttac tccaaaaaga tggacccagg    60
197 tccgaagggg cactgccact gtggggggca tggccatcct ccaggtcact gcgggccacc   120
198 ccctggccat ggcccagggc cctgcgggcc accccccac catgggccag ggccctgcgg    180
199 gccacccccc caccatggtc caggccctg cgggccaccc cctggccatg gcccagggcc    240
200 ctgcgggcca ccccccacc atggtccagg ccctgcgggg cctccccctg gccatggccc    300
201 aggtcaccca cccctggtc cacatcactg aggaagtaga agaaaacagg acacaagatg    360
202 gcaagcctga gagaattgcc cagctgacct ggaatgaggc ctaaaccaca atcttctctt   420
203 cctaataaac agcctcctag aggccacatt ctattcttta aaaaaaaaa aaaaaaaaa    480
204 aaaaaaaaa aaaaaaaaa aaaaaaaaa aaaa                                  514
206 <210> SEQ ID NO: 13
207 <211> LENGTH: 1893
208 <212> TYPE: DNA
209 <213> ORGANISM: Homo sapiens
211 <220> FEATURE:
212 <221> NAME/KEY: misc_feature
213 <222> LOCATION: (1184)..(1184)
214 <223> OTHER INFORMATION: n equals a,t,g, or c
216 <220> FEATURE:
217 <221> NAME/KEY: misc_feature
218 <222> LOCATION: (1865)..(1865)
219 <223> OTHER INFORMATION: n equals a,t,g, or c
221 <220> FEATURE:
222 <221> NAME/KEY: misc_feature
223 <222> LOCATION: (1883)..(1883)
224 <223> OTHER INFORMATION: n equals a,t,g, or c
226 <220> FEATURE:
227 <221> NAME/KEY: misc_feature
228 <222> LOCATION: (1887)..(1887)
229 <223> OTHER INFORMATION: n equals a,t,g, or c
231 <220> FEATURE:
232 <221> NAME/KEY: misc_feature
233 <222> LOCATION: (1893)..(1893)
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     234 <223> OTHER INFORMATION: n equals a,t,g, or c
     236 <400> SEQUENCE: 13
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     238 agtagccgcc tctgccgccg cggagcttcc cgaacctctt cagccgcccg gagccgctcc     120
     239 cggagcccgg ccgtagaggc tgcaatcgca gccgggagcc cgcagcccgc gccccgagcc     180
     240 cgccgccgcc cttcgagggc gccccaggcc gcgccatggt gaaggtgacg ttcaactccg     240
     241 ctctggccca gaaggaggcc aagaaggacg agcccaagag cggcgaggag gcgctcatca     300
     242 tcccccccga cgccgtcgcg gtggactgca aggacccaga tgatgtggta ccagttggcc     360
     243 aaagaagagc ctggtgttgg tgcatgtgct ttggactagc atttatgctt gcaggtgtta     420
     244 ttctaggagg agcatacttg tacaaatatt ttgcacttca accagatgac gtgtactact     480
     245 gtggaataaa gtacatcaaa gatgatgtca tcttaaatga gccctctgca gatgccccag     540
     246 ctgctctcta ccagacaatt gaagaaaata ttaaaatctt tgaagaagaa gaagttgaat     600
     247 ttatcagtgt gcctgtccca gagtttgcag atagtgatcc tgccaacatt gttcatgact     660
     248 ttaacaagaa acttacagcc tatttagatc ttaacctgga taagtgctat gtgatccctc     720
     249 tgaacacttc cattgttatg ccacccagaa acctactgga gttacttatt aacatcaagg     780
     250 ctggaaccta tttgcctcag tcctatctga ttcatgagca catggttatt actgatcgca     840
     251 ttgaaaacat tgatcacctg ggtttcttta tttatcgact gtgtcatgac aaggaaactt     900
     252 acaaactgca acgcagagaa actattaaag gtattcagaa acgtgaagcc agcaattgtt     960
     253 tcgcaattcg gcattttgaa aacaaatttg ccgtggaaac tttaatttgt tcttgaacag    1020
     254 tcaagaaaaa cattattgag gaaaattaat atcacagcat aaccccaccc tttacatttt    1080
     255 gtgcagtgat tattttttaa agtcttcttt catgtaagta gcaaacaggg ctttactatc    1140
W--> 256 ttttcatctc attaattcaa ttaaaaccat taccttaaaa tttnaaaaaa aaaaaaaaaa    1200
     257 aggcccgccg cgctcgcctc tccgccccgc gtccagctcg cccagctcgc ccagcgtccg    1260
     258 ccgcgcctcg gccaaggctt caacggacca caccaaaatg ccatctcaaa tggaacacgc    1320
     259 catggaaacc atgatgttta catttcacaa attcgctggg gataaaggct acttaacaaa    1380
     260 ggaggacctg agagtactca tggaaaagga gttccctgga tttttggaaa atcaaaaaga    1440
     261 ccctctggct gtggacaaaa taatgaagga cctggaccag tgtagagatg gcaaagtggg    1500
     262 cttccagagc ttcttttccc taattgcggg cctcaccatt gcatgcaatg actattttgt    1560
     263 agtacacatg aagcagaagg gaaagaagta ggcagaaatg agcagttcgc tcctccctga    1620
     264 taagagttgt cccaaagggt cgcttaagga atctgcccca cagcttcccc catagaagga    1680
     265 tttcatgagc agatcaggac acttagcaaa tgtaaaaata aaatctaact ctcatttgac    1740
     266 aagcagagaa agaaaagtta aataccagat aagcttttga tttttgtatt gtttgcatcc    1800
     267 ccttgccctc aataaataaa gttctttttt agttccaaaa aaaaaaaaag ggcggccgtt    1860
W--> 268 taarngatcc aasttacgta ccntgcntgc gan                                 1893
     270 <210> SEQ ID NO: 14
     271 <211> LENGTH: 1681
     272 <212> TYPE: DNA
     273 <213> ORGANISM: Homo sapiens
     275 <400> SEQUENCE: 14
     276 gcccgctgag gacgcagcgt cagctgacct ggggagtcgc gattcgtgcc ggccggtcct      60
     277 ggttctccgg tcccgccgct cccgcagcag ccatgtcgtt cttcccggag ctttacttta     120
     278 acgtggacaa tggctacttg gagggactgg tgcgcggcct gaaggccggg gtgctcagcc     180
     279 aggccgacta cctcaacctg gtgcagtgcg agacgctaga ggacttgaaa ctgcatctgc     240
     280 agagcactga ttatggtaac ttcctggcca acgaggcatc acctctgacg gtgtcagtca     300
     281 tcgatgaccg gctcaaggag aagatggtgg tggagttccg ccacatgagg aaccatgcct     360
     282 atgagccact cgccagcttc ctagacttca ttacttacag ttacatgatc gacaacgtga     420
     283 tcctgctcat cacaggcacg ctgcaccagc gctccatcgc tgagctcgtg cccaagtgcc     480
     284 acccactagg cagcttcgag cagatggagg ccgtgaacat tgctcagaca cctgctgagc     540
     285 tctacaatgc cattctggtg gacacgcctc ttgcggcttt tttccaggac tgcatttcag     600
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 08/09/2002
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Input Set : A:\PA003P1.seqList.txt
Output Set: N:\CRF4\08092002\I997003.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 3
Seq#:13; N Pos. 1184,1865,1883,1887,1893
Seq#:24; N Pos. 812,822,829,838,841
Seq#:25; N Pos. 329,332
Seq#:27; N Pos. 738,771
Seq#:28; N Pos. 1014
Seq#:29; N Pos. 419
Seq#:46; Xaa Pos. 215,216,227,236
Seq#:47; Xaa Pos. 153

VERIFICATION SUMMARY   DATE: 08/09/2002
PATENT APPLICATION: US/09/997,003   TIME: 14:56:29

Input Set : **A:\PA003Pl.seqList.txt**
Output Set: **N:\CRF4\08092002\I997003.raw**

```
L:7 M:270 C: Current Application Number differs, Replaced Current Application No
L:7 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:256 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1140
L:268 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1860
L:545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:780
L:546 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:840
L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:300
L:615 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:720
L:659 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:960
L:686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:360
L:1305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 after pos.:208
L:1308 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 after pos.:224
L:1351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:144
```